

1600

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DATE: 07/24/2002
                RAW SEQUENCE LISTING
                                                       TIME: 12:08:33
                PATENT APPLICATION: US/09/522,727D
                Input Set : A:\PTO.AMC.txt
                Output Set: N:\CRF3\07242002\I522727D.raw
 4 <110> APPLICANT: DANA-FARBER CANCER INSTITUTE, INC.
        MARASCO, Wayne
        MHASHILKAR, Abner
 8 <120> TITLE OF INVENTION: INTRABODY-MEDIATED CONTROL OF IMMUNE REACTIONS
10 <130> FILE REFERENCE: 47577 C
12 <140> CURRENT APPLICATION NUMBER: 09/522,727D
13 <141> CURRENT FILING DATE: 2000-03-10
15 <150> PRIOR APPLICATION NUMBER: PCT/US98/19563
16 <151> PRIOR FILING DATE: 1998-09-18
18 <150> PRIOR APPLICATION NUMBER: 60/059,339
19 <151> PRIOR FILING DATE: 1997-09-19
21 <160> NUMBER OF SEQ ID NOS: 56
23 <170> SOFTWARE: PatentIn version 3.1
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 15
27 <212> TYPE: PRT
28 <213> ORGANISM: human
30 <400> SEQUENCE: 1
31 Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Ser
32 1
34 <210> SEQ ID NO: 2
35 <211> LENGTH: 15
36 <212> TYPE: PRT
37 <213> ORGANISM: human
39 <400> SEQUENCE: 2
40 Glu Ser Gly Arg Ser Gly Gly Gly Gly Ser Gly Gly Gly Ser
41 1
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43 <210> SEQ ID NO: 3
44 <211> LENGTH: 14
45 <212> TYPE: PRT
46 <213> ORGANISM: human
48 <400> SEQUENCE: 3
49 Glu Gly Lys Ser Ser Gly Ser Gly Ser Glu Ser Lys Ser Thr
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52 <210> SEO ID NO: 4
53 <211> LENGTH: 15
54 <212> TYPE: PRT
55 <213> ORGANISM: human
57 <400> SEQUENCE: 4
58 Glu Gly Lys Ser Ser Gly Ser Gly Ser Glu Ser Lys Ser Thr Gln
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61 <210> SEQ ID NO: 5
62 <211> LENGTH: 14
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63 <212> TYPE: PRT
64 <213> ORGANISM: human
66 <400> SEQUENCE: 5
67 Glu Gly Lys Ser Ser Gly Ser Gly Ser Glu Ser Lys Val Asp
68
   1
                     5
70 <210> SEQ ID NO: 6
71 <211> LENGTH: 14
72 <212> TYPE: PRT
73 <213> ORGANISM: human
75 <400> SEQUENCE: 6
76 Gly Ser Thr Ser Gly Ser Gly Lys Ser Ser Glu Gly Lys Gly
77
                     5
79 <210> SEQ ID NO: 7
80 <211> LENGTH: 18
81 <212> TYPE: PRT
82 <213> ORGANISM: human
84 <400> SEQUENCE: 7
85 Lys Glu Ser Gly Ser Val Ser Ser Glu Gln Leu Ala Gln Phe Arg Ser
                     5
86
   1
                                        10
87 Leu Asp
90 <210> SEQ ID NO: 8
91 <211> LENGTH: 16
92 <212> TYPE: PRT
93 <213> ORGANISM: human
95 <400> SEQUENCE: 8
96 Glu Ser Gly Ser Val Ser Ser Glu Glu Leu Ala Phe Arg Ser Leu Asp
99 <210> SEQ ID NO: 9
100 <211> LENGTH: 35
101 <212> TYPE: DNA
102 <213> ORGANISM: human
104 <400> SEQUENCE: 9
105 tttgcggccg ctcaggtgca rctgctcgag tcygg
                                                                             35
107 <210> SEQ ID NO: 10
108 <211> LENGTH: 66
109 <212> TYPE: DNA
110 <213> ORGANISM: human
112 <400> SEOUENCE: 10
113 agatecgeeg ceaeegetee caccacetee ggagecaeeg ceaeetgagg tgaeegtgae
                                                                             60
114 crkggt
                                                                             66
116 <210> SEQ ID NO: 11
117 <211> LENGTH: 69
118 <212> TYPE: DNA
119 <213> ORGANISM: human
121 <400> SEQUENCE: 11
122 ggtggcggtg gctccggagg tggtgggagc ggtggcggcg gatctgagct cswgmtgacc
                                                                             60
123 cagtctcca
                                                                             69
125 <210> SEQ ID NO: 12
126 <211> LENGTH: 47
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127 <212> TYPE: DNA
128 <213> ORGANISM: human
130 <400> SEQUENCE: 12
131 gggtctagac tcgaggatcc ttattaacgc gttggtgcag ccacagt
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133 <210> SEQ ID NO: 13
134 <211> LENGTH: 6
135 <212> TYPE: PRT
136 <213> ORGANISM: human
138 <400> SEQUENCE: 13
139 Ser Glu Lys Asp Glu Leu
140
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142 <210> SEQ ID NO: 14
143 <211> LENGTH: 59
144 <212> TYPE: DNA
145 <213> ORGANISM: human
147 <400> SEQUENCE: 14
148 gggtctagac tcgaggatcc ttattacagc tcgtcctttt cgcttggtgc agccacagt
                                                                             59
150 <210> SEQ ID NO: 15
151 <211> LENGTH: 24
152 <212> TYPE: DNA
153 <213> ORGANISM: human
155 <400> SEQUENCE: 15
156 tttaccatgg aacatctgtg gttc
                                                                             24
158 <210> SEQ ID NO: 16
159 <211> LENGTH: 30
160 <212> TYPE: DNA
161 <213> ORGANISM: human
163 <400> SEQUENCE: 16
164 ttagcgcgct gaggtgaccg tgaccrkggt
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166 <210> SEQ ID NO: 17
167 <211> LENGTH: 4
168 <212> TYPE: PRT
169 <213> ORGANISM: human
171 <400> SEQUENCE: 17
172 Lys Asp Glu Leu
173
    1
175 <210> SEO ID NO: 18
176 <211> LENGTH: 4
177 <212> TYPE: PRT
178 <213> ORGANISM: human
180 <400> SEQUENCE: 18
181 Asp Asp Glu Leu
182
184 <210> SEQ ID NO: 19
185 <211> LENGTH: 4
186 <212> TYPE: PRT
187 <213> ORGANISM: human
189 <400> SEQUENCE: 19
190 Asp Glu Glu Leu
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193 <210> SEQ ID NO: 20
194 <211> LENGTH: 4
195 <212> TYPE: PRT
196 <213> ORGANISM: human
198 <400> SEQUENCE: 20
199 Gln Glu Asp Leu
200
    1
202 <210> SEQ ID NO: 21
203 <211> LENGTH: 4
204 <212> TYPE: PRT
205 <213> ORGANISM: human
207 <400> SEQUENCE: 21
208 Arg Asp Glu Leu
209
211 <210> SEO ID NO: 22
212 <211> LENGTH: 7
213 <212> TYPE: PRT
214 <213> ORGANISM: human
216 <400> SEQUENCE: 22
217 Pro Lys Lys Lys Arg Lys Val
218 1
220 <210> SEQ ID NO: 23
221 <211> LENGTH: 7
222 <212> TYPE: PRT
223 <213> ORGANISM: human
225 <400> SEQUENCE: 23
226 Pro Gln Lys Lys Ile Lys Ser
227 1
229 <210> SEQ ID NO: 24
230 <211> LENGTH: 5
231 <212> TYPE: PRT
232 <213> ORGANISM: human
234 <400> SEQUENCE: 24
235 Gln Pro Lys Lys Pro
236 1
238 <210> SEO ID NO: 25
239 <211> LENGTH: 12
240 <212> TYPE: PRT
241 <213> ORGANISM: human
243 <400> SEQUENCE: 25
244 Arg Lys Lys Arg Arg Gln Arg Arg Arg Ala His Gln
247 <210> SEQ ID NO: 26
248 <211> LENGTH: 16
249 <212> TYPE: PRT
250 <213> ORGANISM: human
252 <400> SEQUENCE: 26
253 Arg Gln Ala Arg Arg Asn Arg Arg Arg Trp Arg Glu Arg Gln Arg
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15
     254
           1
                                              10
     256 <210> SEQ ID NO: 27
     257 <211> LENGTH: 19
     258 <212> TYPE: PRT
     259 <213> ORGANISM: human
     261 <400> SEQUENCE: 27
     262 Met Pro Leu Thr Arg Arg Pro Ala Ala Ser Gln Ala Leu Ala Pro
     263
     264 Pro Thr Pro
     267 <210> SEQ ID NO: 28
     268 <211> LENGTH: 15
     269 <212> TYPE: PRT
     270 <213> ORGANISM: human
     272 <400> SEQUENCE: 28
     273 Met Asp Asp Gln Arg Asp Leu Ile Ser Asn Asn Glu Gln Leu Pro
     276 <210> SEQ ID NO: 29
     277 <211> LENGTH: 32
     278 <212> TYPE: PRT
     279 <213> ORGANISM: human
     281 <220> FEATURE:
     282 <221> NAME/KEY: UNSURE
     283 <222> LOCATION: (7)(8)(32)
     284 <223> OTHER INFORMATION: UNSURE
     286 <400> SEQUENCE: 29
W--> 287 Met Leu Phe Asn Leu Arg Xaa Xaa Leu Asn Asn Ala Ala Phe Arg His
     288
                           5
W--> 289 Gly His Asn Phe Met Val Arg Asn Phe Arg Cys Gly Gln Pro Leu Xaa
     290
                                          25
     292 <210> SEQ ID NO: 30
     293 <211> LENGTH: 8
     294 <212> TYPE: PRT
     295 <213> ORGANISM: human
     297 <400> SEQUENCE: 30
     298 Gly Cys Val Cys Ser Ser Asn Pro
     299
         1
     301 <210> SEQ ID NO: 31
     302 <211> LENGTH: 8
     303 <212> TYPE: PRT
     304 <213> ORGANISM: human
     306 <400> SEQUENCE: 31
     307 Gly Gln Thr Val Thr Thr Pro Leu
         1
     310 <210> SEQ ID NO: 32
     311 <211> LENGTH: 8
     312 <212> TYPE: PRT
     313 <213> ORGANISM: human
     315 <400> SEQUENCE: 32
     316 Gly Gln Glu Leu Ser Gln His Glu
```

RAW SEQUENCE LISTING ERROR SUMMARY
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the $\langle 220 \rangle$ to $\langle 223 \rangle$ fields of each sequence which presents at least one n or Xaa.

Seq#:29; Xaa Pos. 7,8,32

Seq#:51; N Pos. 505
Seq#:51; Xaa Pos. 169
Seq#:52; Xaa Pos. 169